SCORE Search Results Details for Application 10 Result 20090127_104823_us-10-682-1

This page gives you Search Results detail for the Application 10682179 and Search Result 2009012

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: January 27, 2009, 10:51:49 ; Search time 2016 Seconds
(without alignments)
74969.960 Million cell updates/sec

Title: US-10-682-179A-1
Perfect score: 1495
Sequence: 1 tgatcatggctcagattgaa.....atgactggggtgaagtcgta 1495

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 14594359 seqs, 50548307366 residues

Total number of hits satisfying chosen parameters: 29188718

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : GenEmbl:*
1: gb_env:*
2: gb_pat:*
3: gb_phg:*
4: gb_pln:*
5: gb_pri:*
6: gb_rod:*
7: gb_sts:*
8: gb_syn:*
9: gb_una:*
10: gb_vrl:*
11: gb_vrt:*
12: gb_inv:*
13: gb_mam:*
14: gb_bct:*
15: gb_htg1:*
16: gb_htg2:*
17: gb_htg3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	% Query